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The potential role of myocardial serotonin receptor 2B expression in canine dilated cardiomyopathy

Fonfara, S ; Hetzel, U ; Oyama, M A ; Kipar, A

Abstract: Serotonin signalling in the heart is mediated by receptor subtype 2B (5-HTR2B). A contribution of serotonin to valvular disease has been reported, but myocardial expression of 5-HTR2B and its role in canine dilated cardiomyopathy (DCM) is not known. The aim of the present study was to investigate myocardial 5-HTR2B mRNA expression in dogs with DCM and to correlate results with expression of markers for inflammation and remodelling. Myocardial samples from eight healthy dogs, four dogs with DCM, five with cardiac diseases other than DCM and six with systemic non-cardiac diseases were investigated for 5-HTR2B mRNA expression using quantitative PCR (qPCR). The results were compared to mRNA expression of selected cytokines, matrix metalloproteinases (MMP) and tissue inhibitors of matrix metalloproteinase (TIMP). Laser microdissection with subsequent qPCR and immunohistochemistry were employed to identify the cells expressing 5-HTR2B. The myocardium of control dogs showed constitutive 5-HTR2B mRNA expression. In dogs with DCM, 5-HTR2B mRNA values were significantly greater than in all other groups, with highest levels of expression in the left ventricle and right atrium. Myocytes were identified as the source of 5-HTR2B mRNA and protein. A significant positive correlation of 5-HTR2B mRNA with expression of several cytokines, MMPs and TIMPs was observed. The findings suggest that serotonin might play a role in normal cardiac structure and function and could contribute to myocardial remodelling and functional impairment in dogs with DCM.

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Abstract: Serotonin signalling in the heart is mediated by receptor subtype 2B (5-HTR2B). A contribution of serotonin to valvular disease is reported, but myocardial expression of 5-HTR2B and its role in canine dilated cardiomyopathy (DCM) is not known. The aim of this study was to investigate myocardial 5-HTR2B mRNA expression in dogs with DCM and to correlate results with expression of markers for inflammation and remodelling. Myocardial samples from eight healthy dogs, four dogs with DCM, five dogs with cardiac diseases other than DCM and six dogs with systemic non-cardiac diseases were investigated for 5-HTR2B mRNA expression using quantitative PCR (qPCR). The results were compared to mRNA expression of selected cytokines, matrix metalloproteinases (MMP) and tissue inhibitors of matrix metalloproteinase (TIMP). Laser microdissection with subsequent qPCR and immunohistochemistry were employed to identify the cells expressing 5-HTR2B. The myocardium of control dogs showed constitutive 5-HTR2B mRNA expression. In dogs with DCM, 5-HTR2B mRNA values were significantly greater than in all other groups, with highest levels of expression in the left ventricle and right atrium. Myocytes were identified as the source of 5-HTR2B mRNA and protein. A significant positive correlation of 5-HTR2B with expression of several cytokines, MMPs and TIMPs was observed. These findings suggest that serotonin might play a role in normal cardiac structure and function and could contribute to myocardial remodelling and functional impairment in dogs with DCM.

Dear Brian,

I did proof-read the manuscript and have addressed the additional queries. I did use track-changes to mark what I have changed/added and for comments. I did check the references and all the references in the text are present on the reference list and vice versa.

Best regards

Sonja

Original Article

The potential role of myocardial serotonin 2B receptor expression in canine dilated cardiomyopathy

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29 **Abstract**

30 Serotonin signalling in the heart is mediated by receptor subtype 2B (5-HTR2B). A
31 contribution of serotonin to valvular disease is reported, but myocardial expression of 5-
32 HTR2B and its role in canine dilated cardiomyopathy (DCM) is not known. The aim of this
33 study was to investigate myocardial 5-HTR2B mRNA expression in dogs with DCM and to
34 correlate results with expression of markers for inflammation and remodelling. Myocardial
35 samples from eight healthy dogs, four dogs with DCM, five dogs with cardiac diseases other
36 than DCM and six dogs with systemic non-cardiac diseases were investigated for 5-HTR2B
37 mRNA expression using quantitative PCR (qPCR). The results were compared to mRNA
38 expression of selected cytokines, matrix metalloproteinases (MMP) and tissue inhibitors of
39 matrix metalloproteinase (TIMP). Laser microdissection with subsequent qPCR and
40 immunohistochemistry were employed to identify the cells expressing 5-HTR2B. The
41 myocardium of control dogs showed constitutive 5-HTR2B mRNA expression. In dogs with
42 DCM, 5-HTR2B mRNA values were significantly greater than in all other groups, with
43 highest levels of expression in the left ventricle and right atrium. Myocytes were identified as
44 the source of 5-HTR2B mRNA and protein. A significant positive correlation of 5-HTR2B
45 with expression of several cytokines, MMPs and TIMPs was observed. These findings
46 suggest that serotonin might play a role in normal cardiac structure and function and could
47 contribute to myocardial remodelling and functional impairment in dogs with DCM.

48

49 *Keywords:* Serotonin, Heart failure, Cardiac disease, Dilated cardiomyopathy, Myocardial
50 remodelling

51

Introduction

Serotonin (5-hydroxytryptamine; 5-HT) is a monoamine neurotransmitter, which is produced from tryptophan by two different tryptophan hydroxylases (TPH), namely, TPH1 and TPH2, found in enterochromaffine cells of the gastrointestinal tract and in neurons of the central nervous system, respectively (Jonnakuty and Gragnoli, 2008). Serotonin is involved in platelet function, vascular and non vascular smooth muscle contraction as well as cardiac function (Nebigil and Maroteaux, 2003; Jonnakuty and Gragnoli, 2008). It is rapidly removed from the circulation, via cellular uptake by the serotonin transporter (SERT) and subsequently stored in platelets or metabolised in pulmonary vascular endothelial cells and hepatocytes by monoamine oxidase (Jonnakuty and Gragnoli, 2008). However, high circulating 5-HT concentrations or administration of serotonergic drugs are associated with arrhythmia and valvulopathy (Sheline et al., 1997; reviewed by Kaumann and Levy, 2006; Elangbam et al., 2008; Orton et al., 2012). Furthermore, 5-HT is suspected to be involved in myxomatous valvular disease (MVD) in humans and dogs (Fitzgerald et al., 2000; Arndt et al., 2009; Oyama and Levy, 2010).

Serotonin exerts its effect through seven different receptor groups (5-HTR1 to 5-HTR7), composed of several receptor subtypes (Elangbam et al., 2005; Kaumann and Levy, 2006), all of which are members of the G-protein-coupled receptor superfamily. In the cardiovascular system, the receptor subtypes 5-HTR1B, 2A, 2B, 4 and 7 can be found, of which the receptor subtype 5-HTR2B, which is present on endothelial cells, smooth muscle cells, fibroblasts, valvular interstitial cells and cardiomyocytes, is suspected to be involved in cardiac remodelling and development of valvulopathies (Rothman et al., 2000; Kaumann and Levy, 2006; Disatian and Orton, 2009; Oyama and Levy, 2010; Hutcheson et al., 2011).

77 Activation of 5-HTR2B has been found to stimulate phospholipase C and A2, both of
78 which increase the intracellular calcium concentration. In addition, downstream signalling,
79 involving extracellular-signal regulated kinase (ERK), mediates proliferative effects by
80 inducing transcription of transforming growth factor (TGF)- β and other effector genes, such
81 as matrix metalloproteinases (MMP) and bone morphogenic protein, which potentially
82 contribute to the pathogenesis of mitral valve disease (Disatian and Orton, 2009; Lacerda et
83 al., 2012a; Orton et al., 2012). Furthermore, activation of 5-HTR2B has been shown to
84 increase expression of interleukin (IL)-1, IL-6 and tumor necrosis factor alpha (TNF- α)
85 mRNA and protein in murine cardiac fibroblasts (Jaffre et al., 2004). These inflammatory
86 cytokines are known to be elevated in the blood of humans and dogs affected with congestive
87 heart failure (CHF) and are involved in cardiac inflammation and remodelling (Anker and
88 von Haehling, 2004; Fonfara et al., 2011). Tensile strain has been suspected to increase TPH
89 and 5-HTR2B and reduce SERT in canine septal mitral valve leaflets, which might result in
90 local 5-HT synthesis and prolonged 5-HT activity (Elangbam et al., 2008; Scruggs et al.,
91 2010; Lacerda et al., 2012a; Lacerda et al., 2012b).

92
93 The role of 5-HT and its receptors in canine DCM is not fully understood. Dilated
94 cardiomyopathy is associated with left ventricular dilatation and ventricular wall thinning,
95 therefore increasing wall stress. Increased ventricular wall stress is likely to stimulate
96 mechanoreceptors, which could cause local 5-HT production and increased 5-HTR2B
97 expression, as reported for valve leaflets exposed to tensile strain (Disatian and Orton, 2009;
98 Lacerda et al., 2012b). This appears likely, since in rats, chronic 5-HT administration has
99 been shown to increase valvular 5-HTR2B transcription (Elangbam et al., 2008). In humans
100 with heart failure, 5-HT plasma concentrations are also elevated, which has led to the
101 assumption that heart failure is mediated by 5-HTR2B (Jaffre et al., 2009; Shyu, 2009).

102

103 | It is suspected that through induction of IL-1, IL-6, TNF- α and TGF- β 1, 5-HT
104 | contributes to cardiomyocyte hypertrophy, increased fibrosis and ventricular stiffness leading
105 | to reduced cardiac contractility and heart disease (Anker and von Haehling, 2004; Jaffre et
106 | al., 2004; Disatian and Orton, 2009; Jaffre et al., 2009). Increased concentrations of TGF- β 1,
107 | a potent profibrotic cytokine, have been reported in canine MVD (Aupperle et al., 2008;
108 | Disatian and Orton, 2009) and CHF (Fonfara et al., 2011). We have recently shown that dogs
109 | with end-stage cardiac diseases exhibit increased myocardial mRNA expression of
110 | inflammatory cytokines, TGF- β 1, MMP-2 as well as tissue inhibitor of matrix
111 | metalloproteinase (TIMP)-1 and TIMP-2 (Fonfara et al., 2013a; Fonfara et al., 2013b).
112 | Considering these results and the characteristic pathological changes in canine DCM, we
113 | hypothesize that the 5-HT system plays a role in the pathogenesis of canine DCM. We
114 | therefore designed a study to investigate cardiac expression of 5-HTR2B and its association
115 | with expression of IL-1, IL-6, TNF- α , TGF- β 1, MMP-1, -2, -3, -13, TIMP-1 and -2,
116 | comparing healthy control dogs with other groups of dogs affected with DCM, other cardiac
117 | diseases or systemic non-cardiac disease.

118

119 **Material and methods**

120 *Animals and tissues*

121 | Details of clinical cases are shown in Table 1 and consist of dogs affected with DCM
122 | (group 2; $n = 4$), cardiac diseases other than DCM (group 3; $n = 5$) and dogs with systemic,
123 | non-cardiac disease (group 4; $n = 6$) (Fonfara et al., 2013a; Fonfara et al., 2013b). Eight
124 | control Beagles (group 1^a) were used for the quantitative PCR assay (four each entire male
125 | and female, sourced from a pharmaceutical company with a median age of 2.75 years). The
126 | dogs were euthanized and post-mortem examination performed on site. Samples from the

127 myocardium (left and right atrium [LA, RA], left and right ventricle [LV, RV]) were
128 immersed in RNAlater (Ambion) and provided for this project. Two Doberman Pinscher dogs
129 (one female neutered 2 years old, one male entire 6 years old), with no gross or histological
130 evidence of cardiac disease, served as controls for the immunohistological examination
131 (~~group 1b~~).

Comment [11]: I have removed the group, because they are not in one of the groups, we did compare. So in a way they don't need a group and I felt the 1a and 1b was confusing.

133 All of the dogs with cardiac and systemic diseases were clinical cases that had
134 undergone diagnostic investigations according to their underlying disease and presenting
135 clinical signs. Investigations were performed by clinical specialists in their respective fields
136 or board-registered residents under supervision. For cardiac cases, investigations included a
137 cardiac work-up, comprising blood pressure measurement, electrocardiography,
138 echocardiography and thoracic radiography, at different time points prior to death (Fonfara et
139 al., 2013b). Diagnoses were made, based on applying standardised criteria for clinical
140 assessment of cardiac cases. For classification of heart failure, the ABCD scheme was used
141 (Strickland, 2008). Dogs had been euthanized upon owners' request, due to poor prognosis
142 and impaired quality of life, with one exception, where a dog developed ventricular
143 fibrillation and died (Fonfara et al., 2013b). Informed consent was obtained from owners
144 prior to inclusion into the study and samples were anonymised by assigning identification
145 numbers. The study was approved by the University of Bristol Committee on Research
146 Ethics.

148 Hearts were removed within 1 h of death and examined for any gross pathological
149 abnormalities. Myocardial samples (interventricular septum [IVS], RA, RV, LA, LV) were
150 collected and stored in RNAlater at -20 °C until use. Hearts were subsequently fixed in 10%
151 formalin for a minimum of 48 h and tissue samples from the same sites as those for RNA

152 extraction were prepared and paraffin wax embedded, according to routine procedures for
153 histological and immunohistological examinations. The hearts from the immunohistology
154 control dogs (~~group 1b~~) were formalin-fixed and subsequently processed as for the other
155 hearts.

156

157 *Laser microdissection*

158 A left ventricular sample from a dog with DCM stored in RNAlater was embedded in
159 OCT compound (Tissue-Tek) and frozen at -40 °C. Cryosections (8-10 µm) were prepared
160 and placed onto specific membrane slides (Carl Zeiss), which had been incubated in dry heat
161 at 180 °C for 4 h prior to use, in order to inactivate RNase enzymes. The slides with
162 cryosections were air dried, stained with haematoxylin as recommended by the manufacturer
163 (Carl Zeiss) and immediately used.

164

165 Cardiomyocytes were isolated with a Zeiss Palm microbeam laser microdissection
166 microscope, using a 40× lens. Cutting was performed with 52-55% energy, 60-62% focus and
167 10% speed and collection with 65% energy and 47% focus. Three samples of approximately
168 100-150 cardiomyocytes were collected into a 500 µL adhesive cap tube (Carl Zeiss). After
169 collection, tissue was lysed into 50 µL RLT buffer containing 2-mercaptoethanol (Qiagen).

170 Samples were subsequently frozen at -20 °C for 18-22 h and ~~subsequently~~ RNA isolation was
171 performed using the RNeasy Minikit (Qiagen).

172

173 *Quantitative assessment of 5-HTR2B mRNA expression*

174 Total RNA was extracted from myocardial samples and cDNA synthesized as reported
175 previously (Fonfara et al., 2011b). For the samples obtained by laser microdissection, 14 µL
176 RNase-free water was used to elute RNA from the column prior to cDNA synthesis. The

Comment [I2]: Otherwise it is double?

177 ~~forward and reverse~~ ~~Published~~ primer sequences ~~were used~~ for the canine housekeeping
178 gene GAPDH ~~were~~ 5'CTGGGGCTCACTTGAAAGG3' ~~and~~
179 5'CAAACATGGGGGCATCAG3', ~~respectively,~~ ~~and~~ ~~for~~ ~~5-HTR2B~~
180 5'CCCAATGAGGCTCTGCAGTT3' ~~and~~
181 5'CTGTGATGAGAAGTGTATCTAGTAGAATGATT3', ~~respectively,~~ ~~and~~ ~~5-HTR2B~~
182 [Editor comment: might be better to put primer sequences here. You reference three
183 publications for two primer pairs] (~~Oyama and Chittur, 2006; Fonfara et al., 2013a; Fonfara~~
184 ~~et al., 2013b~~). The primer efficiency of the GAPDH primer pair was 100% and of 5-HTR2B
185 97%. PCR was performed and analyzed according to standard protocols, the expression of 5-
186 HTR2B was normalized to GAPDH expression (relative expression) and calculated via the $2^{-\Delta\Delta C_t}$
187 method, as previously reported (Fonfara et al., 2013a; Fonfara et al., 2013b).

188

189 *Statistical analysis and correlation of 5-HTR2B mRNA with cytokine, MMP and TIMP*
190 *expression*

191 For the statistical analysis of the quantitative PCR results, Minitab 16 was used.
192 Following performance of basic descriptive statistics, 5-HTR2B mRNA values were log
193 transformed to improve normality and the model assumptions necessary for parametric
194 analysis. For comparison of different groups and/or cardiac regions, one-way ANOVA tests
195 were employed. Results are displayed as mean and standard deviation.

196

197 To test for a potential correlation between 5-HTR2B and cytokine, MMP or TIMP
198 expression in the hearts of dogs with DCM, the qPCR results for IL-1, IL-6, TNF- α , TGF- β 1,
199 MMP-1, 2, -3, -13, TIMP-1 and TIMP-2, obtained from the same myocardial samples, were
200 used (Fonfara et al., 2013a; Fonfara et al., 2013b). Expression was examined with scatterplots

201 and then tested with the Pearson correlation test. Statistical significance was defined as $P <$
202 0.05.

203

204 *Histology and immunohistochemistry*

205 For the histological examination, 3-5 μm thick sections were prepared from the paraffin
206 wax-embedded tissue blocks and stained with haematoxylin and eosin (HE).
207 Immunohistochemistry for 5-HTR2B was performed on sections from the RA and the RV
208 and/or LV using a murine anti-human 5-HTR2B antibody (clone A72-1; Beckton Dickinson)
209 and the horseradish peroxidase method (HRP mouse kit, Dako), following antigen retrieval
210 by incubation in citrate buffer (pH 6.0) and microwaving (Disatian and Orton, 2009).
211 Consecutive sections were incubated with an isotype-control mouse monoclonal antibody and
212 sections from canine stomach, tissue from a case of suppurative myocarditis and smooth
213 muscle cells present in myocardial arteries served as positive controls.

214

215 **Results**

216 *Pathology and histopathology*

217 The hearts of dogs with systemic diseases and clinical diagnoses of cardiac diseases
218 other than DCM did not exhibit any gross or histological changes, i.e. there was no evidence
219 of degenerative, neoplastic or inflammatory changes (Fonfara et al., 2013b). The hearts of
220 dogs with DCM showed the typical gross changes (cardiomegaly, biventricular dilatation,
221 myocardial eccentric hypertrophy). Histologically, interstitial, subendo- and subepicardial
222 fibrosis, lipomatosis cordis, leukocyte infiltration, and focal cardiomyocyte necrosis was
223 observed, as previously reported (Fonfara et al., 2013b).

224

225 *Canine cardiac 5-HTR2B expression*

Comment [I3]: Should we possibly say that this served as negative control?

Expression of 5-HTR2B mRNA was detected in the myocardium of control dogs in all examined locations (RA, RV, LA, LV), with some individual variation in quantitative values (Fig. 1). In the other groups, 5-HTR2B expression was also consistently observed, including in the IVS, which was not available for examination in control dogs. There was a greater individual variation in the disease groups (Table 2), although for groups 3 and 4, 5-HTR2B values did not differ significantly from those of control dogs. In contrast, dogs with DCM (group 2) demonstrated significantly greater 5-HTR2B mRNA expression when compared to all other groups (Table 2).

The results of the RT-PCR performed on isolated cardiomyocytes of the LV of a dog affected with DCM, showed that the cardiomyocytes were expressing 5-HTR2B mRNA and the immunohistochemical staining of both control and diseased dogs confirmed that cardiomyocytes also synthesised 5-HTR2B protein (Fig. 2). Immunohistochemistry showed 5-HTR2B protein expression not only in cardiomyocytes, but also in vascular smooth muscle cells and infiltrating as well as intravascular neutrophils (Fig. 2). Immunostaining in cardiomyocytes was generally weak and often patchy or with negative fibers intermingled with positive fibers. However, staining in cardiomyocytes was generally most intense in the right atrium, a finding that was supported by the general trend for greatest mRNA expression at this location (Fig. 1).

Disease-associated differences in 5-HTR2B mRNA expression ~~values~~ concentrations and patterns

A comparison of 5-HTR2B mRNA ~~values~~ concentrations in the different groups showed comparable expression levels in controls dogs and those with cardiac diseases other than DCM, or dogs with systemic diseases ($P = 0.46$). However, in dogs with DCM, mRNA

Comment [14]: One of the reviewers did not like 'values' and requested this to be changed to 'concentrations'.

251 ~~values-concentrations~~ were significantly greater than in all other groups ($P = 0.002$; Fig. 1).
252 This difference was primarily due to greater 5-HTR2B mRNA ~~values-concentrations~~ in the
253 LV ($P = 0.005$) and the RA ($P = 0.027$) of dogs with DCM, when the different cardiac
254 regions were compared between groups. In dogs with DCM, the same significant difference
255 in 5-HTR2B mRNA ($P = 0.001$) was observed between the different regions, with 5-HTR2B
256 mRNA ~~concentrations values~~ in the RA also being significantly higher than in the LV ($P =$
257 0.047 ; Fig. 1). None of the other groups exhibited significant differences in mRNA
258 ~~concentrations values~~ comparing different cardiac locations.

259

260 The immunohistochemical results confirmed the mRNA expression data, since staining
261 of cardiomyocytes for 5-HTR2B was more consistent and more intense in the RA, compared
262 to other locations in the DCM dogs, and compared to the same location in dogs from other
263 groups (Fig. 2B, C).

264

265 *Correlation between 5-HTR2B and cytokine, MMP and TIMP expression in the myocardium*
266 *of dogs with DCM*

267 We have previously shown upregulation of cytokine, MMP and TIMP mRNA
268 expression in the myocardium of dogs with DCM (Fonfara et al., 2013a; Fonfara et al.,
269 2013b). Using the qPCR data generated from the same RNA extracts and applying the
270 Pearson's correlation test, a significant positive correlation was seen between 5-HTR2B
271 mRNA expression and IL-1 ($r = 0.719$, $P < 0.001$; Fig. 3a), IL-6 ($r = 0.624$, $P = 0.003$), TNF-
272 α ($r = 0.661$, $P = 0.002$), TGF- β 1 ($r = 0.848$, $P < 0.001$; Fig. 3b), MMP-2 ($r = 0.493$, $P =$
273 0.027), MMP-13 ($r = 0.770$, $P < 0.001$; Fig. 3a), TIMP-1 ($r = 0.729$, $P < 0.001$; Fig. 3b) and
274 TIMP-2 ($r = 0.826$, $P < 0.001$). [Editor comment: should you add an additional figure to
275 illustrate this data ? What are the r values ?]

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276

277 Discussion

278 The present study assesses the constitutive expression of 5-HTR2B in the myocardium
279 of dogs and the changes in expression pattern and intensity in association with canine DCM.
280 The examination of control dogs showed that the myocardium constitutively expresses 5-
281 HTR2B, suggesting a role for 5-HT and its cardiac receptor in normal cardiac structure and
282 function (Nebigil et al., 2001; Nebigil et al., 2003; Liang et al., 2006). There were no
283 significant differences in regional expression of 5-HTR2B mRNA, but a trend for higher
284 expression in the RA was observed.

Comment [15]: You have removed this result from the results section, should we still keep this comment in the discussion?

285

286 Our results indicate that severe systemic diseases and non-DCM cardiac diseases do not
287 influence 5-HTR2B expression, whereas canine DCM is associated with a significant
288 increase in the RA and, to a lesser, but still significant extent, the LV. This 5-HTR2B
289 upregulation correlated with increased IL-1, IL-6, TNF- α , TGF- β 1, MMP-2, MMP-13,
290 TIMP-1 and TIMP-2 mRNA expression. Using laser microdissection and immunohistology,
291 we were able to confirm that cardiomyocytes synthesize 5-HTR2B, which is similar to
292 previously reported results for canine MVD and cardiomyocytes in mice and humans (Choi
293 and Maroteaux, 1996; Kaumann and Levy, 2006; Liang et al., 2006; Disatian and Orton,
294 2009; Oyama and Levy, 2010; Orton et al., 2012).

295

296 Serotonin, in association with increased 5-HTR2B and reduced SERT production, has
297 been reported in dogs with MVD (Oyama and Chittur, 2006; Arndt et al., 2009; Disatian and
298 Orton, 2009; Scruggs et al., 2010). The results of the present study indicate that 5-HTR2B
299 also plays a role in canine DCM. In contrast to the results of the current study, in mouse
300 models of DCM, reduced 5-HTR2B protein expression was observed, whereas increased 5-

301 HTR2B expression caused cardiac hypertrophy (Nebigil et al., 2001; Nebigil et al., 2003).
302 Since the dogs with DCM in our study exhibited ventricular eccentric hypertrophy and were
303 all in decompensated CHF, these results may not be entirely contradictory, but are on the
304 other hand not fully comparable.

305

306 In humans, CHF has been shown to be associated with an increase of 5-HT
307 independently of the type of cardiac disease (Jaffre et al., 2009). Since chronic 5-HT
308 administration results in increased 5-HTR2B expression (Elangbam et al., 2008), it is
309 possible that 5-HT is also chronically elevated in dogs that develop DCM, resulting in
310 increased 5-HTR2B transcription, thereby suggesting that CHF and not the underlying
311 cardiac disease is responsible for the increase of 5-HT and its receptor. However, if high 5-
312 HT concentrations would be the cause of 5-HTR2B upregulation in DCM, it can be expected
313 that all regions investigated and not just the RA and LV would increase 5-HTR2B gene
314 expression. Instead, our results suggested local changes in the 5-HT system, and upregulation
315 of 5-HTR2B in the LV (which is mainly affected in canine DCM) further supports this
316 hypothesis.

317

318 Interestingly in humans, administration of a 5-HT re-uptake inhibitor has been reported
319 to cause DCM and cardiogenic shock, but myocardial 5-HTR2B was not investigated in this
320 patient (Charniot et al., 2010). It is also possible that an increase of TPH1 and phosphorylated
321 ERK, or a reduction of SERT would result in enhanced serotonin production and signalling,
322 as reported for valvular diseases (Elangbam et al., 2008; Disatian and Orton, 2009; Scruggs et
323 al., 2010; Lacerda et al., 2012a; Lacerda et al., 2012b). Further investigations, in particular
324 assessment of circulating serotonin concentrations and the myocardial signal transduction
325 pathway, would be needed to confirm this. The lack of evidence of increased 5-HTR2B gene

326 expression in dogs with cardiac diseases other than DCM might be a consequence of the case
327 selection, i.e. dogs with a variety of cardiac diseases other than DCM were included into this
328 group.

329

330 We identified a positive correlation between 5-HTR2B expression and IL-1, IL-6, TNF-
331 α , TGF- β 1, MMP-2, [MMP-13](#), TIMP-1 and TIMP-2 mRNA in the myocardium of dogs with
332 DCM. Others have reported that 5-HT can induce IL-1, IL-6, TNF- α , TGF- β 1, MMP-1,
333 MMP-3 and MMP-13 (Jaffre et al., 2004; Yabanoglu et al., 2009; Lacerda et al., 2012b). The
334 inflammatory cytokines IL-1, IL-6 and TNF- α are important mediators of myocardial
335 inflammation and affect myocardial contractility (Anker and von Haehling, 2004). The
336 cytokine TGF- β 1 mediates growth of cardiomyocytes (Schultz et al., 2002) and is suspected
337 to impair mitochondrial energy metabolism (Huntgeburth et al., 2011) and might therefore be
338 involved in eccentric cardiac hypertrophy and contribute to systolic dysfunction in dogs with
339 DCM and CHF. Furthermore, myofibroblast activation by TGF- β 1 alongside mechanical
340 strain is suspected to lead to increased stiffening of heart valves (Orton et al., 2012), and
341 could also be involved in increased ventricular stiffness and functional impairment in dogs
342 with DCM and CHF.

343

344 Matrix metalloproteinases and TIMPs are important regulators of cardiac remodeling
345 (Spinale, 2007). It has been reported that 5-HT induces MMP-3 and MMP-13 expression in
346 cardiac fibroblasts (Yabanoglu et al., 2009) and an increased MMP-1 and MMP-13 protein
347 production in mitral valve leaflets exposed to tensile strain was suspected to be caused by 5-
348 HTR2B stimulation (Lacerda et al., 2012a; Lacerda et al., 2012b). In the present study, a
349 significant positive correlation of 5-HTR2B with MMP-2, MMP-13, TIMP-1 and TIMP-2
350 expression was detected in myocardial samples of dogs with DCM and CHF. Matrix

351 metalloproteinase 2, a gelatinase, and MMP-13, a collagenase, might be involved in matrix
352 degradation and ventricular dilatation, whereas TIMP-1 and -2 might contribute to increased
353 fibrosis and ventricular stiffness in these dogs (Spinale, 2007).

354

355 Increased 5-HTR2B mRNA expression in the RA and the further increase in dogs with
356 DCM is of interest. In humans, elevated 5-HT concentrations have been reported to cause
357 pathological changes in the right heart valves, whereas the left heart valves are presumed to
358 be protected by 5-HT breakdown by pulmonary monoamine oxidase (Hutcheson et al., 2011).
359 Additionally, increased 5-HT concentrations have been shown to produce arrhythmia,
360 including atrial fibrillation (AF) (Sheline et al., 1997). An increase of 5-HTR2B in the RA
361 might reflect enhanced 5-HT signalling. This would contribute to the structural remodelling
362 of the RA, which is an important factor in the development of AF (Brundel et al., 2005).
363 Also, we recently reported increased expression of IL-1, TNF- α and TGF- β 1 in the RA and of
364 MMP-2 and MMP-13 in both atria of dogs with AF (Fonfara et al., 2013a; Fonfara et al.,
365 2013b). TGF- β 1 is known to modulate atrial ion channels, potential promoting AF (Ramos-
366 Mondragon et al., 2011) and MMP-2 and -13 might cause atrial dilatation, which is
367 frequently associated with AF. Therefore, increased 5-HTR2B expression in the RA, together
368 with upregulation of inflammatory cytokines, TGF- β 1, MMP-2 and -13 might contribute to
369 development of AF, a clinical feature observed in three of the four dogs with DCM.

370

371 Arrhythmia might also be caused by 5-HTR2B-induced activation of phospholipase C
372 and A2, which results in an increase in intracellular calcium concentration and disrupted
373 calcium handling (Hutcheson et al., 2011). This might be one of the mechanisms underlying
374 reduced myocardial function observed in dogs with DCM and CHF (Martin et al., 2009;
375 Oyama et al., 2009).

376

377 The study has a number of limitations, including the small number of dogs used, the
378 heterogeneity of groups 3 and 4 and the age difference between the control animals and those
379 in the other groups. Larger sample sizes and groups with more homogeneous phenotypes
380 would have allowed a comparison of different cardiac diseases and might have provided
381 more significant results. Investigation of 5-HTR2B expression in isolation, without studying
382 further components of the signal transduction pathway, limits interpretation of the study
383 findings, in terms of how receptor upregulation might impact on cellular responses. However,
384 a positive correlation of 5-HTR2B expression with increased expression of the downstream
385 gene products of the signal transduction pathway was present. Only dogs with end-stage
386 diseases were included, which also excluded investigation of the involvement of 5-HTR2B in
387 progression of DCM.

388

389 **Conclusions**

390 Constitutive expression of 5-HTR2B mRNA in myocardial samples of young Beagle
391 dogs suggests an involvement of 5-HT and its receptor in normal cardiac structure and
392 function. Increased 5-HTR2B mRNA expression and a positive correlation with IL-1, IL-6,
393 | TNF- α , TGF- β 1, MMP-2, MMP-13, TIMP-1 and TIMP-2 in dogs with DCM and CHF
394 suggests a contribution of 5-HT to structural myocardial remodelling and functional
395 impairment in these dogs. Further investigations of the role of 5-HT, its receptor and the
396 signal transduction pathway in the progression of canine DCM are needed in particular to
397 investigate any potential therapeutic implications.

398

399 **Conflict of interest statement**

None of the authors of this paper has a financial or personal relationship with other people or organizations that could inappropriately influence or bias the content of the paper.

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582
583

584 **Table 1.** Details of dogs in each of the study groups.

Study group	Breed	Age (years)	Sex	Disease	CHIEF classification	BW (kg)	AF (Y/N)
2	Doberman	10	NF	Dilated cardiomyopathy	D	34.8	Y
2	Doberman	8	F	Dilated cardiomyopathy	D	39.6	Y
2	Bullmastiff	7	M	Dilated cardiomyopathy	D	54	N
2	Great Dane	6	M	Dilated cardiomyopathy	D	70.8	Y
3	Boxer	11	M	Aortic stenosis	B	26	N
3	Labrador	0.2	M	Tricuspid dysplasia	C3	7.8	N
3	Labrador	9	NF	Pulmonic stenosis	B	32.5	N
3	Labrador	6	NM	Arrhythmogenic cardiomyopathy	C3	32	N
3	German Shepherd	14	NF	Myxomatous valvular disease	D	22.7	Y
4	Boxer	7.5	M	Brain tumour	-	37.5	N
4	Cocker	8	NF	Brain tumour	-	14.4	N
4	Cross	7	NF	Lymphoma	-	9.6	N
4	Cross	11	NF	Pancreatic carcinoma	-	20.5	N
4	Rottweiler	0.75	M	Secondary hyperparathyroidism	-	24.55	N
4	Labrador	0.5	M	Spinal fracture	-	27.3	N

585

586 CHIEF, Canine Heart Failure International Expert Forum; AF, atrial fibrillation; BW, body

587 weight; CHF, congestive heart failure; F, female; M, male; N, no; NF, neutered female; NM,

588 neutered male; Y, yes.

590 **Table 2.** Relative 5-HTRB2 mRNA expression in different myocardial regions of control
 591 dogs (group 1), dogs with dilated cardiomyopathy (group 2), those with cardiac diseases
 592 other than dilated cardiomyopathy (group 3) and those with systemic non-cardiac diseases
 593 (group 4). Results are displayed as mean (top value) and standard deviation (bottom value).
 594

Region	Group 1 (n = 8)	Group 2 (n = 4)	Group 3 (n = 5)	Group 4 (n = 6)
Interventricular septum	n.d.	1.64 (0.44) ¹	1.37 (0.46)	1.34 (0.50)
Left atrium	1.39 (0.37)	1.62 (0.26) ¹	1.40 (0.33)	1.32 (0.49)
Left ventricle	1.47 (0.20) ²	2.12 (0.26) ^{1,2}	1.20 (0.36) ²	1.22 (0.55) ²
Right atrium	1.97 (0.42) ³	2.89 (0.55) ^{1,3}	1.52 (0.90) ³	2.02 (0.57) ³
Right ventricle	1.27 (0.54)	1.54 (0.34) ¹	1.46 (0.46)	1.08 (0.57)
All samples	1.54 (0.45) ⁴	1.96 (0.62) ⁴	1.39 (0.60) ⁴	1.40 (0.60) ⁴

595

596 **Significant difference between results:** ¹ $P = 0.001$, ² $P = 0.005$, ³ $P = 0.027$, ⁴ $P = 0.002$.

597 n.d., not determined.

598 ¹ A significant difference in 5-HTRB2 mRNA expression was present comparing cardiac
 599 regions of group 2 ($P = 0.001$).

600 ^{2,3} 5-HTRB2 mRNA expression was greater in the left ventricle (² $P = 0.005$) and right atrium
 601 (³ $P = 0.027$) of dogs from group 2, comparing these regions between groups.

602 ⁴ A significant difference in 5-HTRB2 mRNA expression was present comparing all samples
 603 of the different groups ($P = 0.002$).

604 [Editor comment: what does # represent? Can you be more precise about what the statistics
 605 represent?]

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Comment [I6]: Was from the first version where we had signs instead of numbers.

Figure Legends

Fig. 1. Relative 5-HTR2B mRNA expression (normalized to GAPDH expression) in myocardial samples of different cardiac regions [interventricular septum (IVS), left atrium (LA), left ventricle (LV), right atrium (RA) right ventricle (RV)] comparing control dogs (group 1, $n = 8$), dogs with dilated cardiomyopathy (group 2, $n = 4$), dogs with cardiac diseases other than dilated cardiomyopathy (group 3, $n = 5$) and dogs with systemic non-cardiac diseases (group 4, $n = 6$). Significantly greater 5-HTR2B mRNA expression was present in left ventricular and right atrial samples of dogs with DCM ($P < 0.05$). Individual value plot with mean \oplus and median \otimes . [Editor comment: Is the data normally distributed (I assume not) in which case it might be better just to have a median line. You need a y-axis label.]

Fig. 2. Immunohistochemical analysis of 5-HTR2B protein expression in the myocardium of the right atrium. A) Control dog. Cardiomyocytes exhibiting a weak positive cytoplasmic reaction for 5-HTR2B (arrows) are found close to negative cardiomyocytes (arrowheads). B) Dog with DCM. All cardiomyocytes exhibit moderate cytoplasmic staining with an intensity similar to that seen in smooth muscle cells of small arterial walls (arrow). C) Dog with severe pulmonary stenosis. Bundles of weakly positive cardiomyocytes are seen adjacent to aggregates of negative cardiomyocytes (arrowheads). Arrow: small artery. D) Dog with brain tumour. Cardiomyocytes exhibit a weak cytoplasmic reaction. Arrow: small artery. Horseradish peroxidase method, Papanicolaou's haematoxylin counterstain. Bar = 20 μm .

Comment [17]: The results were normalised for analysis (line 192-104). I added both, because that was suggested by the pathologists, who gave feedback after my presentation. I can remove one, but in that case possibly better the mean?

630 **Fig. 3.** Scatter plots showing a positive correlation between 5-HTR2B mRNA expression and
631 *IL-1* ($r = 0.719$, $P < 0.001$), *MMP-13* ($r = 0.770$, $P < 0.001$) (3a), *TGF- β* ($r = 0.848$, $P <$
632 0.001) and *TIMP-1* ($r = 0.729$, $P < 0.001$) (3b) mRNA concentrations.

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633 Editor comment: is it worthwhile illustrating the correlation data, comparing 5-HTRB2 with
634 cytokine, MMP and TIMP values.

Comment [18]: I did select the ones, which had similar values and could therefore be displayed in the same diagrams.

Figure 1

Figure 1

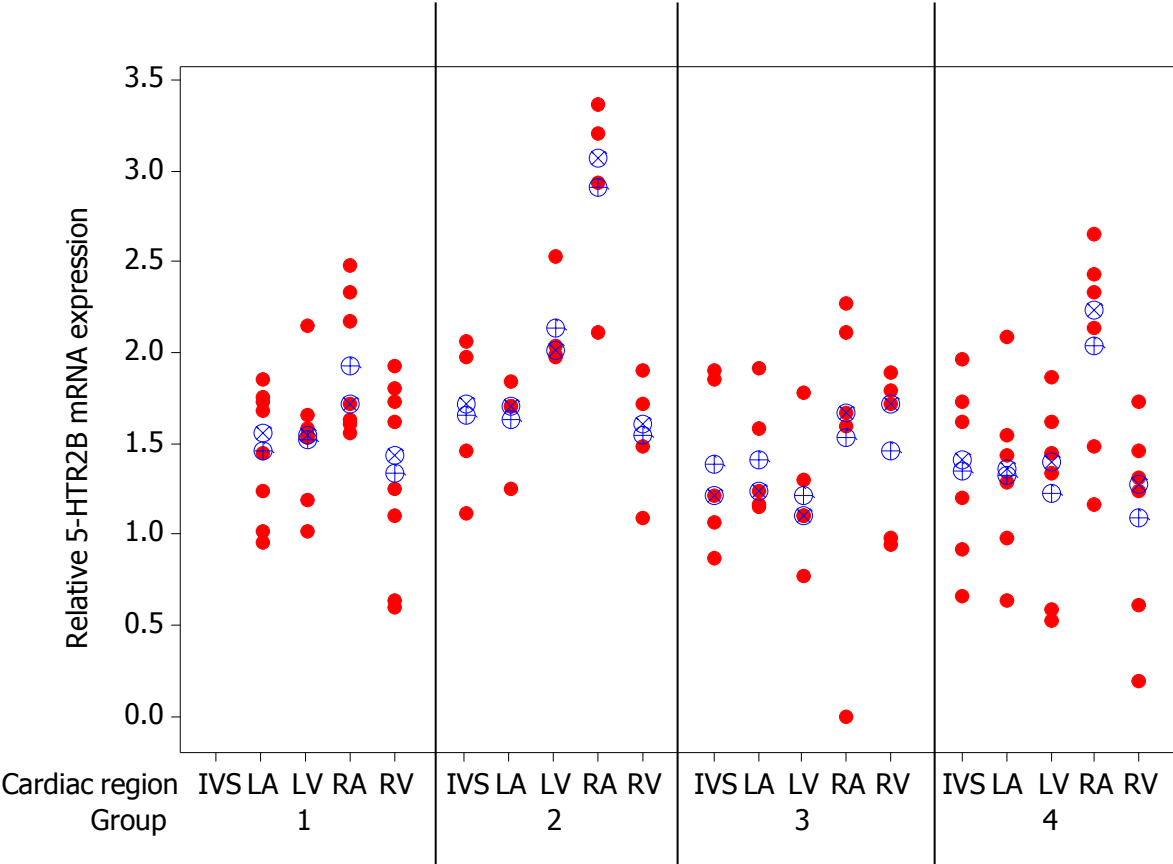


Figure 2
[Click here to download high resolution image](#)

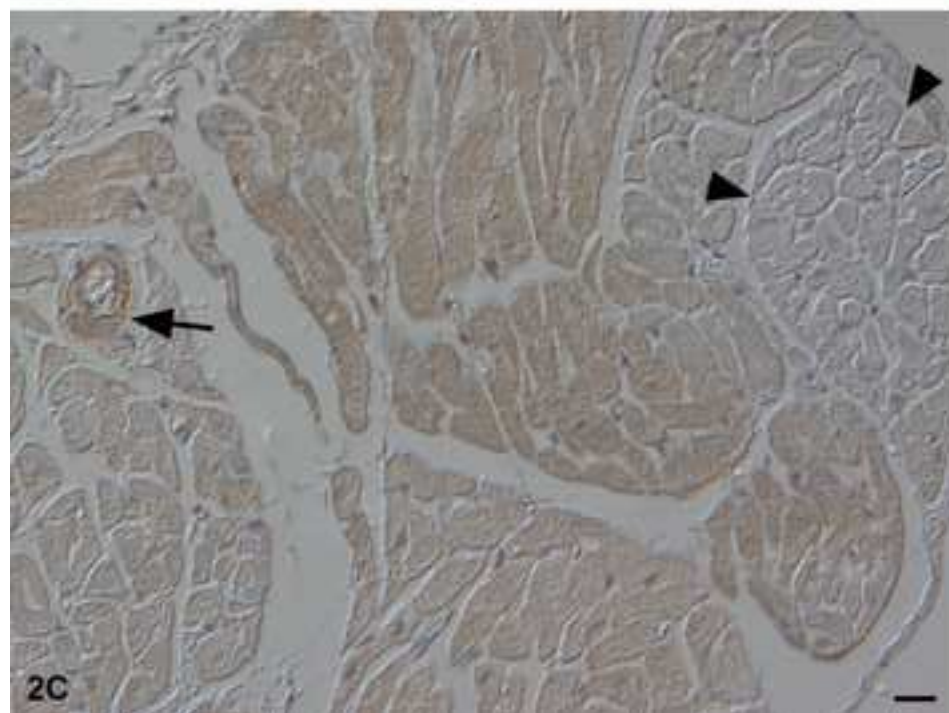
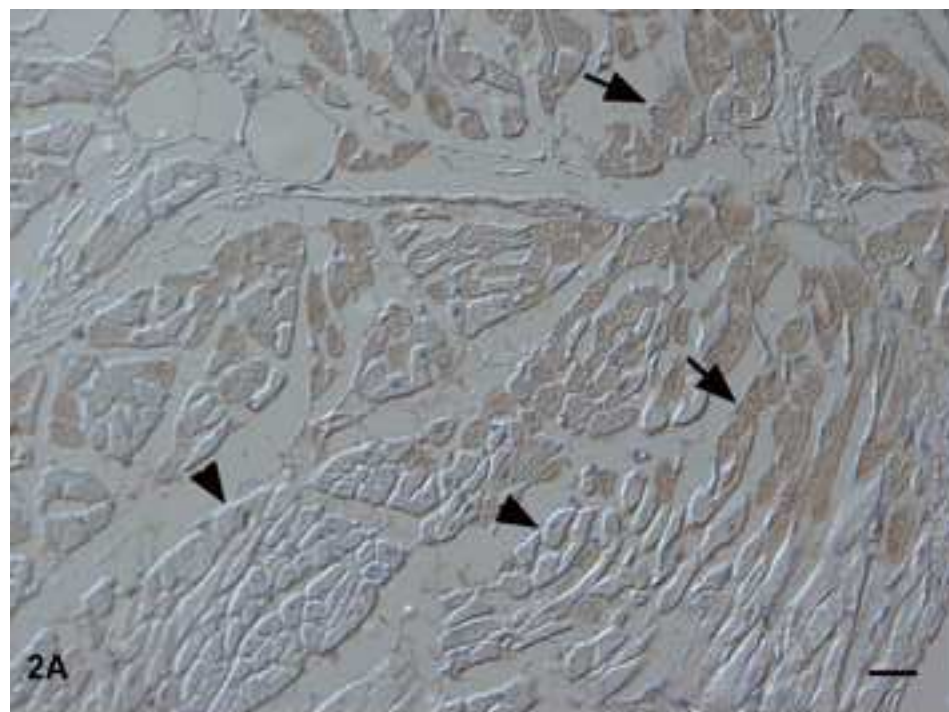


Figure 3

Figure 3a

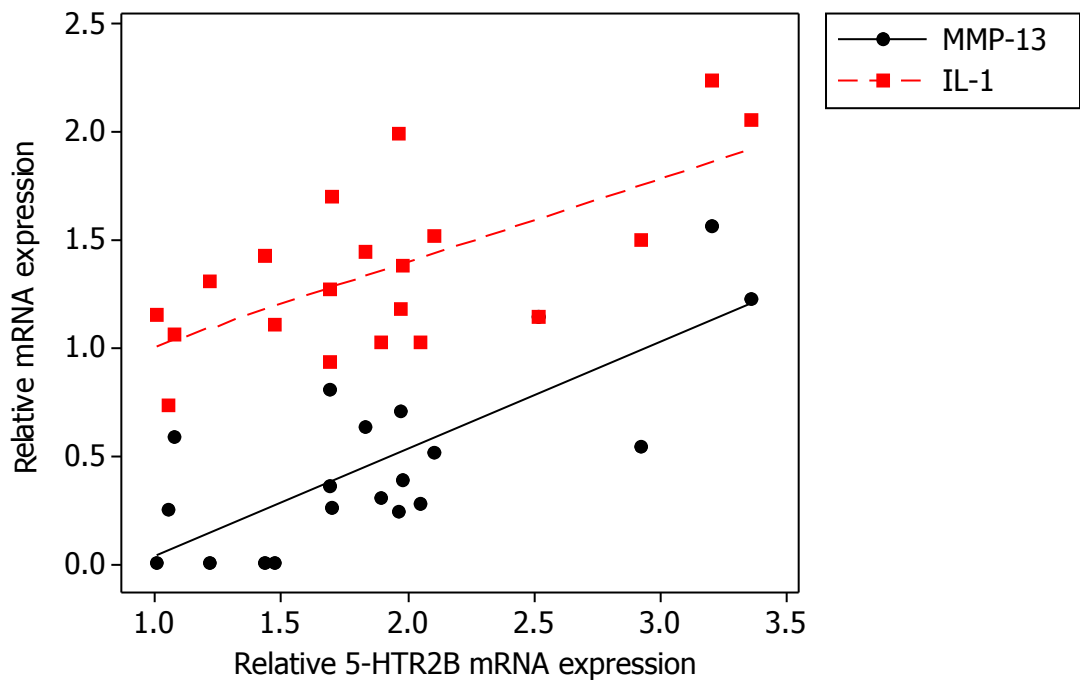


Figure 3b

